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Alignment View Print

	1 10 20 30
seq1	C C T T C T C C	G C C C T T C C C	C T C G C T T T G
seq25	- - - - -	- - - - -	- - - - -
Consensus	[C C T T C T C C]	[G C C C T T C C C]	[C T C G C T T T G]
	40	50	60
seq1	C C T [G G] C G C	T C C T [C C T C C]	C [C] C [C] C C C
seq25	- - - - -	- - - - -	- - - - -
Consensus	[C C T [G G] C G C]	[T C C T [C C T C C]]	[C [C] C [C] C C C]
	70	80	90
seq1	A C C T [G T] G T C C	C C C [G C G C C C G]	C T C C [C C C G]
seq25	- - - - -	- - - - -	- - - - -
Consensus	[A C C T [G T] G T C C]	[C C C [G C G C C C G]]	[C T C C [C C C G]]
	100	110	120
seq1	C [G G] C C T G A G	C C C C T C T C T G	C T [G C C] G C C
seq25	- - - - -	- - - - -	- - - - -
Consensus	[C [G G] C C T G A G]	[C C C C T C T C T G]	[C T [G C C] G C C]
	130	140	150
seq1	C C C C C T [C C T G]	C C C [C T C T C C]	T [G C T] C T C T C G G
seq25	- - - - -	- - - - -	- - - - -
Consensus	[C C C C C T [C C T G]]	[C C C [C T C T C C]]	[T [G C T] C T C T C G G]
	160	170	180
seq1	G T T C T G A G G C	A C [G C T T G T C]	A C C C G G G C
seq25	- - - - -	- - - - -	- - - - -
Consensus	[G T T C T G A G G C]	[A C [G C T T G T C]]	[A C C C G G G C]
	190	200	210
seq1	[G G] A T T C T C T T	T C T C T T T C T C	T T C T [G G] C C C
seq25	- - - - -	- - - - -	- - - - -
Consensus	[[G G] A T T C T C T T]	[T C T C T T T C T C]	[T T C T [G G] C C C]

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Alignment View Print

	430		440		450
seq1					
seq25					
Consensus					
	460		470		480
seq1					
seq25					
Consensus					
	490		500		510
seq1					
seq25					
Consensus					
	520		530		540
seq1					
seq25					
Consensus					
	550		560		570
seq1					
seq25					
Consensus					
	580		590		600
seq1					
seq25					
Consensus					
	610		620		630
seq1					
seq25					
Consensus					

Alignment View Print

	640	650	660
seq1	C C G G . . .	C C C C T T T	C C T G C C C
seq25	C G G G . . .	C C C G T T G T	C T T G C C C
Consensus	C C G G . . .	C C C C T T G T	C C T G C C C
	670	6	690
seq1	C G G C C C T T	C T G C T C G T	G G G
seq25	C G G C C C T T	C T G C T C G T	G G G
Consensus	C G G C C C T T	C T G C T C G T	G G G
	700	710	720
seq1	C C C T G C C G C	C C C G T G C C G C	C C T T C G C C C
seq25	C C C T G C C G C	C C C G T G C C G C	C C T T C G C C C
Consensus	C C C T G C C G C	C C C G T G C C G C	C C T T C G C C C
	730	740	750
seq1	C C T C C G C C C	G G G C C G C G	C T G C G G G
seq25	C C T C C G C C C	G G G C C G C G	C T G C G G G
Consensus	C C T C C G C C C	G G G C C G C G	C T G C G G G
	760	770	780
seq1	G G G C C C G G	G G T C G G C C	C C C T G T G T C
seq25	G G G C C C G G	G G T C G G C C	C C C T G T G T C
Consensus	G G G C C C G G	G G T C G G C C	C C C T G T G T C
	790	800	810
seq1	G C T G C C C	C C C G G G G C C	T T C T C T C C C
seq25	G C T G C C C	C C C G G G G C C	T T C T C T C C C
Consensus	G C T G C C C	C C C G G G G C C	T T C T C T C C C
	820	830	840
seq1	T G G G C C C T	G G G G T G T	C G C C C C G
seq25	T G G G G C C C T	G G G G T G T	C G C C C C G
Consensus	T G G G G C C C T	G G G G T G T	C G C C C C G

Alignment View Print

	850	860	870
seq1	C C G T C C G	C T G G C T G G T G	C G G C C C C G
seq25	C C G T C C G	C T G G C T G G T G	C G G C C C C G
Consensus	C C G T C C G	C T G G C T G G T G	C G G C C C C G
	880	890	900
seq1	G G C T G G C C	C G C G C T C C C	C C T G G G T T T
seq25	G G C T G G C C	C G C G C T C C C	C C T G G G T T T
Consensus	G G C T G G C C	C G C G C T C C C	C C T G G G T T T
	910	920	930
seq1	G G T G G T T T C T	C T C G G G G C	C T C G T C T C G
seq25	G G T G G T T T C T	C T C G G G G C	C T C G T C T C G
Consensus	G G T G G T T T C T	C T C G G G G C	C T C G T C T C G
	940	950	960
seq1	T C T T G T T T G	C T C C C G T T	G G C C T T C
seq25	T C T T G T T T G	C T C C C G T T	G G C C T T C
Consensus	T C T T G T T T G	C T C C C G T T	G G C C T T C
	970	980	990
seq1	T T G T G T G	A G G G G T G	C C G G G G T G
seq25	T T G T G T G	A G G G G T G	C C G G G G T G
Consensus	T T G T G T G	A G G G G T G	C C G G G G T G
	1000	1010	1020
seq1	T G T G T C A	G G T G T C G	T C C G T C C G
seq25	T G T G T C A	G G T G T C G	T C C G T C C G
Consensus	T G T G T C A	G G T G T C G	T C C G T C C G
	1030	1040	1050
seq1	G G G G C C	G G G G C G	G G G T C G G C C
seq25	G G G G C C	G G G G C G	G G G T C G G C C
Consensus	G G G G C C	G G G G C G	G G G T C G G C C

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	1060	1070	1080
seq1	C G T C T T G	G C C C T C G	G C C C C T C C G
seq25	C G T C T T G	G C C C T C G	G C C C C T C C G
Consensus	C G T C T T G	G C C C T C G	G C C C C T C C G
	1090	1100	1110
seq1	C G T C C C C	G G T G G C C T G	G G G G C
seq25	C G T C C C C	G G T G G C C T G	G G G G C
Consensus	C G T C C C C	G G T G G C C T G	G G G G C
	1120	1130	1140
seq1	T C C C T C T T	C C G G G G G	G C C C C C
seq25	T C C C T C T T	C C G G G G G	G C C C C C
Consensus	T C C C T C T T	C C G G G G G	G C C C C C
	1150	1160	1170
seq1	A C T G C C C C	A G C T C T G C	C C C C G C C C
seq25	A C T G C C C C	A G C T C T G C	C C C C G C C C
Consensus	A C T G C C C C	A G C T C T G C	C C C C G C C C
	1180	1190	1200
seq1	A G G T C C T	G C C C C G	C T G C T G
seq25	A G G T C C T	G C C C C G	C T G C T G
Consensus	A G G T C C T	G C C C C G	C T G C T G
	1210	1220	1230
seq1	G G C T G T C C	C C T G C G	C C C C G G C
seq25	G G C T G T C C	C C T G C G	C C C C G G C
Consensus	G G C T G T C C	C C T G C G	C C C C G G C
	1240	1250	1260
seq1	C C G G G C S T T	G G G G G G C T C C G	C C C T G G G C T G
seq25	C C G G G C S T T	G G G G G G C T C C G	C C C T G G G C T G
Consensus	C C G G G C S T T	G G G G G G C T C C G	C C C T G G G C T G

Alignment View Print

	1270	1280	1290
seq1 C T T C C G T C T	C C T C C . . T G C C G G C G T
seq25			
ConsensusCTTCCGTC.....	CC.....TGC.....GGCGT.....
	1300	1310	1320
seq1 G G G C C C C T G	C T G G G C G T G C	G C T G G G C C C C
seq25			
ConsensusGGGCCCCGT.....	CTGGGCGT.....	GCGGGCCCC.....
	1330	1340	1350
seq1 C C T C G T G C C C	T T C C C T T G C C C	C C T G T C C G G C
seq25			
ConsensusCCCTCGT.....	TTCCC.....	CC.....GGC.....
	1360	1370	1380
seq1 C C T G G G G C C T	C T G T T C T G C T	G T G G C C T G C
seq25			
ConsensusCCGGGGCCT.....	CTGTTCTGCT.....	G.....GGCGCTG.....
	1390	1400	1410
seq1 C T C C C C G G G	T C C T G G G G G	C G G C C C C G T
seq25			
ConsensusCTCCCGGG.....	TCC.....GGGGG.....	CG.....GGCGGT.....
	1420	1430	1440
seq1 T G C C C C T C G C	T C A C G C C C C	C C C C C C C G C
seq25			
ConsensusTGCCCCCTCG.....	TCA.....GCCCG.....	CC.....CCCCCG.....
	1450	1460	1470
seq1 C C T C C T G G G C C	C G G C C C C G G G	C G G C C C T T C G
seq25			
ConsensusCTCCGGCC.....	CG.....GCCCGG.....	CG.....CCCTTCG.....

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	1480	1490	1500
seq1	CCC C [C] C T [G]	T C T [G] C [C] C [G] T C	T [C] C T C T T [G] T
seq25			
Consensus	[C] C C C [C] C T [G]	T C T [G] C [C] C [G] T C	T [C] C T C T T [G] T
	1510	1520	1530
seq1	[G] G C C T C [G] C	[G] G C C T C [G] G C C C	C [G] G G C C T [G] C
seq25			
Consensus	[G] G C C T C [G] C	[G] G C C T C [G] G C C C	C [G] G G C C T [G] C
	1540	1550	1560
seq1	C T C [C] C [G] C C	A [G] G C T G A [C] T	[G] G G C T T [G] G C T G
seq25			
Consensus	C T C [C] C [G] C C	A [G] G C T G A [C] T	[G] G G C T T [G] G C T G
	1570	1580	1590
seq1	C [C] T G T C C T G	T T T [G] T G G C T	C C [C] C T C G G
seq25			
Consensus	C [C] T G T C C T G	T T T [G] T G G C T	C C [C] C T C G G
	1600	1610	1620
seq1	G T G A T T T T	C T T T T T G C C A	T T T G T T T C C
seq25			
Consensus	G T G A T T T T	C T T T T T G C C A	T T T G T T T C C
	1630	1640	1650
seq1	G @ T C C T G T C T	T C T T T T G T C	A T G C C T C G T
seq25			
Consensus	G @ T C C T G T C T	T C T T T T G T C	A T G C C T C G T
	1660	1670	1680
seq1	G T A T T T G G G G	G T G C T G T G T G	G G G G G C T G T
seq25			
Consensus	G T A T T T G G G G	G T G C T G T G T G	G G G G G C T G T

Alignment View Print

